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(74) Agents: CASIMIR, David, A. et al.; Casimir Jones, 440 Science Drive. Suite 203. Madison. WI 53711 (US). (10) International Publication Number WO 2008/104002 A3

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(54) Title: METHODS FOR RAPID FORENSIC DNA ANALYSIS

67) Abstract: The present invention provides methods and primer pairs for rapid, high-resolution forensic analysis of DNA and STR typing by using amplification and mass septermentry, determining the molecular masses and calculating base compositions of amplification products and comparing the molecular masses with the molecular masses of theoretical amplicons indexed in a database.

International application No PCT/US2008/054926

Relevant to claim No.

1-10,17,

A. CLASSIFICATION OF SUBJECT MATTER INV. C12Q1/68

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols) C12Q

Category* | Citation of document, with indication, where appropriate, of the relevant passages

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EPO-Internal, EMBASE, BIOSIS, WPI Data

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Date of the	actual completion of like international search	Date of mailing of the international sea	irch report	
2	9 August 2008	26/01/2009		
Name and	mailing address of the ISA/ European Patent Office, P.B. 5818 Palentiaan 2 NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Fax: (+31-70) 340-3016	Authorized officer Hennard, Christop	he	

International application No PCT/US2008/054926

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International application No. PCT/US2008/054926

Box No. II Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)
This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:
Claims Nos.: because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
Claims Nos.: bocause they are dependent claims and are not drafted in accordance with the second and third centences of Rule 6.4(a).
Box No. III Observations where unity of invention is lacking (Continuation of item 3 of first sheet)
This International Searching Authority found multiple inventions in this international application, as follows:
see additional sheet
 As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. As all searchable claims could be searched without effort justifying an additional fees, this Authority did not Invite payment of additional fees.
 As only some of the required additional search ties were timely paid by the applicant, this international search reportioners only times claims for which fees were pick, specifically deline Nos.
4. \[\] No required additional search bees were Immely gold by the applicant Consequently, this international search report is restricted to the invention first manifold in the claims, it is covered by claims Noc. 1-11, 17, 62-78, 97-106, 110-113, 115 (in part); 12, 30-33 85-86 (entirely) Remark on Protest The additional control free were accomprained by the applicant's protest and, where applicable, the payment of a protest rec.
lee was not paid within the time timit specified in the invitation. No protest accompanied the payment of additional search fees.

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

This International Searching Authority found multiple (groups of) inventions in this international application, as follows:

Invention 1: Claims 1-11, 17, 62-78, 97-106, 110-113, 115 (in part)
and 12, 30-33, 85-86 (entirely)

Concern a method for STR typing comportsing amplifying a nucleic acid using primers, determining the molecular mass of at least one amplification product by mass spectrometry and comparing the molecular mass to a database for matching to the STR allele; an oligonucleotide primer pair; a kit comprising such primer pair in particular wherein the method is typing the STR locus D55818 and further in particular wherein the method is typing the STR locus D55818 and further in particular wherein the method is typing a STR allele which comprises at least one SMP.

Inventions 2-13: Claims 1-11, 13-29, 34-84, 87-106, 110-115 (in part)

Concern a method for STR typing comprising amplifying a nucleic acid using primers, determining the molecular mass to a least one amplification product by mass spectrometry and comparing the molecular mass to a database for matching to the STR allele, an oligonucleotide primer pair; a kit comprising such primer pair wherein, the method is typing the STR locus D8S21179, VMA, D18317, D78820, THO1, TPOX, CSF1PO, D16S539, FGA, D2IS11, D18S51 or D3S respectively of the inventions.

Invention 14: Claims 107-109 entirely

Concern a kit for forensic analysis comprising at least two oligonucleotide primer pairs, one pair configured to hybridise to a region flanking a variable region comprising WKR and the second pair configured to hybridise within or flanking an AMEL locus.

Information on patent family members

International application No PCT/US2008/054926

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